CL

CRF Errors Corrected by the STIC Systems Branch

Serial Number: _08/ 166,925_

CRF Processing Date: _04/28/94_
Edited by: _____ (STIC staff)
Verified by: _____

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: _____

ENTERED

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Inserted a space between the last nucleic designator and the nucleic number for sequences: _____

☐ Deleted page numbers in the text of the sequence listing, which is considered invalid text.

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☐ Deleted non-ASCII "garbage" at the end of files, and other invalid text, such as a secretary's initials. _____

☐ Inserted mandatory headings, specifically: _____

☑ Corrected an obvious error in the response, specifically: _to "(B)Type: nucleic acid"_ _d'ed all "(B)Type:cDNA", to "(B)Type: nucleic acid"_

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Other: _____

_____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

8/01/93

CL

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION   US/08/166,925

DATE: 04/25/94
TIME: 13:33:43

INPUT SET: S7859.raw

This Raw Listing contains only the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

1
2
3   (1)   General Information:
4
5   (i) APPLICANT: Falck-Pedersen, Erik S.
6
7   (ii) TITLE OF INVENTION: ADENOVIRUS GENE EXPRESSION SYSTEM
8
9   (iii) NUMBER OF SEQUENCES: 1
10
11   (iv) CORRESPONDENCE ADDRESS:
12   (A) ADDRESSEE: Alan S. Korman
13   (B) STREET: 1600 Main Place Tower
14   (C) CITY: Buffalo
15   (D) STATE: New York
16   (E) COUNTRY: U.S.A.
17   (F) ZIP: 14202
18
19   (v) COMPUTER READABLE FORM:
20   (A) MEDIUM TYPE: Floppy disk
21   (B) COMPUTER: IBM PC compatible
22   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
23   (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
24
25   (vi) CURRENT APPLICATION DATA:
26   (A) APPLICATION NUMBER: 08/166,925
27   (B) FILING DATE: 12/14/93
28   (C) CLASSIFICATION:
29
30   (viii) ATTORNEY/AGENT INFORMATION:
31   (A) NAME: Korman, Alan S.
32   (B) REGISTRATION NUMBER: 33,932
33   (C) REFERENCE/DOCKET NUMBER: 19603/230
34
35   (ix) TELECOMMUNICATION INFORMATION:
36   (A) TELEPHONE: 716-853-8104
37   (B) TELEFAX: 716-853-8109
38
39
40   (2) INFORMATION FOR SEQ ID NO:1:
41
42   (i) SEQUENCE CHARACTERISTICS:
43   (A) LENGTH: 6783 base pairs
44   (B) TYPE: nucleic acid
45   (C) STRANDEDNESS: double
46   (D) TOPOLOGY: linear

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48      (ii) MOLECULE TYPE: cDNA (genomic)
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52      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION   US/08/166,925

INPUT SET: S7859.raw

Original Text

Line   Error

SEQUENCE LISTING

(1) GENERAL INFORMATION:

    (i) APPLICANT: Falck-Pedersen, Erik S.

    (ii) TITLE OF INVENTION: ADENOVIRUS GENE EXPRESSION SYSTEM

    (iii) NUMBER OF SEQUENCES: 1

    (iv) CORRESPONDENCE ADDRESS:
        (A) ADDRESSEE: Alan S. Korman
        (B) STREET: 1600 Main Place Tower
        (C) CITY: Buffalo
        (D) STATE: New York
        (E) COUNTRY: U.S.A.
        (F) ZIP: 14202

    (v) COMPUTER READABLE FORM:
        (A) MEDIUM TYPE: Floppy disk
        (B) COMPUTER: IBM PC compatible
        (C) OPERATING SYSTEM: PC-DOS/MS-DOS
        (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

    (vi) CURRENT APPLICATION DATA:
        (A) APPLICATION NUMBER: 08/166,925
        (B) FILING DATE: 12/14/93
        (C) CLASSIFICATION:

    (viii) ATTORNEY/AGENT INFORMATION:
        (A) NAME: Korman, Alan S.
        (B) REGISTRATION NUMBER: 33,932
        (C) REFERENCE/DOCKET NUMBER: 19603/230

    (ix) TELECOMMUNICATION INFORMATION:
        (A) TELEPHONE: 716-853-8104
        (B) TELEFAX: 716-853-8109

(2) INFORMATION FOR SEQ ID NO:1:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 6783 base pairs
        (B) TYPE: cDNA   *Change CW1 to "nucleic acid"*
        (C) STRANDEDNESS: double
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: cDNA (genomic)

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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AGAGCTGCAG CTGCCGTCAT CCCTGAGCAG GGGGGCCACT TCGTTAAGCA TGTCCCTGAC 3720

TCGCATGTTT TCCCTGACCA AATCCGCCAG AAGGCGCTCG CCGCCCAGCG ATAGCAGTTC 3780

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CCCGTTCAGC CCGACCGCTG CGCCTTATCC GGTAACTATC G
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